OIPE



ENTERED



TECH CENTER 1600/2900 DATE: 05/05/200

TIME: 07:36:18

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/834,271A

Input Set: N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\I834271A.raw

```
1 4110 APPLICANT: Widner, William
              Sloma, Alan
              Thomas, Michael E.
      4 - 120 - TITLE OF INVENTION: Methods For Producing A polypeptide In a
              Badillus Cell
      → <130 - FILE REFERENCE: 5455.200-US
      " 4140 - CURRENT APPLICATION NUMBER: 09/834,271A
      3 (141 - CURRENT FILING DATE: 2001-04-12
      + 0150 - PRIOR APPLICATION NUMBER: US/09/258,377
     1 | <151 | PHIOR FILING DATE: 1999-02-26
     11 <150 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/031,442
W--> 12 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-26
     13 4160 NUMBER OF SEQ ID NOS: 33
     14 - 1770 FORTWARE: FastSEQ for Windows Version 3.0
     16 :210 SEQ ID NO: 1
     10 000120 IMPE: DNA
19 000130 000BANISM: Bacillus
     20 H4011 SEQUENCE: 1
             aattgelett aagggeeegg gaegteaage ttategatge ggateegegg eege
                                                                                        54
     25 KM1 DE SEQ DE NO: 2
     24 ROMATH LENGTH: 51
     25 HOLDE TYPE: DNA
     26 -0.1130 (#ERANISM: Bacillus
     2" -44000 SEQUENCE: 2
              rosgaatise egggeeetge agttegaata getaegeeta ggegeeggeg e
                                                                                        51
     3 -0.1100 SEQ 11 NO: 3
     51 40110 DENGTH: 58
     3. KU12H TYPE: DNA
     bo KM13: CREANISM: Bacillus
     34 KGCD SEQUENCE: 3
             -agotangest taagggooog ggaogtogag otcaagettg oggoogcoat ggtogaog
     \pm 0.0140 SEQ ID NO: 4
     % -0.11. BENGTH: 58
     39 H. 125 TYPE: DNA
     10 00.130 OFGANISM: Bacillus
     41 -:400: SEQUENCE: 4
     41
              tocqqaatto cogggeootg cagetegagt tegaacgeeg geggtaccag etgettaa.
     44 -0.100 SEQ ID NO: 5
     45 -0111: LENGTH: 37
     46 HOLD TYPE: DNA
     47 (213) CFGANISM: Bacillus
```

48 <400: SEQUENCE: 5

RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/09/834,271A TIME: C7:36:18

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\1834271A.raw

49		ctccgggccc atctgagctc	tataaaaatg	aggaggg		37
		SEO ID NO: 6				
5.2	×11 ·	1ENGTH: 17				
5 3	-1.11.	IYE: DUR				
5.;	11	OF GANISM: Facillus				
515	4000	SEQUENCE: €				
Ę, į,		idfojqatdo atadadaaaa	aaacgct			27
£, -	$\leq 1.0100 \pm$	SEQ ID ID: 7				
£, , •	11111	LEMGTH: 37				
6:	A11	ITTE: DNA				
61	$\{1,1\}, s\in$	CF. WANTEM: Facillus				
6.	1.1	SELUEDOR: 7				
6 -		omaggictta agggoogdat	gegteettet	ttgtgct		37
		3EQ 10 16: 8				
6	-1-11	LEDGTH: 10				
6	-1.11	TYPE: DUA				
(:::-	4.150	CR WANTEM: Earillus				
6,11	-(4:H:	SELUENCE: 8				
10		chagadotho titicaatgig	taacatatga			30
		SE, ID NO: 9	-			
		LENGTH: 42				
		This: DNA				
i, t	·	OBJANISM: Bacillus				
		SEQUEDOE: 9				
		ttiggoritta agggootgca	atogattgtt	tgagaaaaga	aq	42
·, •		SFO ID NO: 10	, ,	3 9 9		
		LEDGTH: 43				
		TYPE: DMA				
		OF WANTEM: Facillus				
		SEQUENCE: 10				
- 1		ttreader.co attitottat	acaaattata	ttttacatat	caq	43
		SBQ ID NO: 11				
		LENGTH: 30				
		TYFE: DNA				
		OFCANISM: Pacillus				
		SECUENCE: 11				
		garaccopy: agotttcagt	daadtacdtd			30
		SEC ID NO: 12	3443444949			
		LED STH: 13				
		TYEE: DNA				
		OEGANISM: Pacillus				
		SEQUENCH: 12				
96		gddyddiae aattcaaag				19
-		· Ship ID NO: 13				
		· LENGTH: 27				
		TYFE: DNA				
		· GRGANISM: Bacillus				
		· SEQUENCE: 13				
105		gggccctcga aacgtaagat	daaacct			27
100		gggccccga aacgcaagac	. gauace			~ /

RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/09/834,271A TIME: 07:36:18

Imput Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\1834271A.raw

109 02	10> SEG ID NO: 14 11> LEUGTH: 29					
	12 TYTE: DNA 13 OPHANISM: Bacillus					
	ON - SEQUENCE: 14					
1:.	parata atabataatt	ttcaaactg				29
	1: - SE, TD NO: 15					
	11 LENGTH: 31					
	10 TYPE: ONA					
	ls OFGANISM: Bacillus OG SERUENCE: 15					
	casponen. To	c				21
	1 - SF, ID No: 16					7.5.2
	11 - Lebsth: 23					
1.13 -0.1	1. · TYPE: DNA					
	l 🐖 OFGANISM: Badillus 🧪					
	On - SEQUENCE: 16					
	paintato titaattaag	ctt				23
	le - SF, 10 ND: 17 11 - DEDGTH: 13					
	1. · TYPE: DNA.					
	ls - OF FAMUSM: Bacillus					
	O - SEQUENCE: 17					
	ga:chuquac thgttcatgt	gaa				23
	le SPO ID NO: Li					
	11 · DENGTH: 23					
	1. TYPE: DNA					
	l - EFGANISM: Badillus Do ESEQUENCE: 18					
1.:	daudtestsa tadataattt	t.ca				23
14	188 SEQ 210 NO: 19					
145 %	110 18073CH: 44					
	l TYFE: INA					
	ls: OF::ANTSN: Bacillus					
	Des SEQUENCE: 19			-+-+		1.7
	gasatusang qqqqttgaca 10: SKG II: NG: EK	ttattttact	gataigiala	alal		44
	11: LFNGTH: 48					
	ILD TYPE: DNA					
	ls: OFGANISM: Bacillus					
15.5	DOTE SEQUENCE: 20					
1: -:	autaavatga ctatacatat	tatattaaac	atattcttt	acctcgag		48
	100 SEQ ID NO: 21					
	11: LENGTH: 3050					
	lib TYFE: INA 180: CEGANISM: a:Badillus					
	DO: SEQUENCE: 21					
161	togaaacgta agatgaaacc	ttagataaaa	gtgcttttt	tgttgcaatt	gaagaattat	60
162	taatgttaag cttaattaaa					120
		-		•		

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/834,271A

DATE: 08/05/2003
TIME: 07:36:19

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\I834271A.raw

163			_	atgtttgaac			120
164				ctaactatag			5:1
16°	-			actigiteat			देशाः
160			-	tacattgtta			កស៊ី()
16	ottgatagos	ttactatacc	tascatgatg	tagtattasa	tgäatatgta	aatatatta	4
165	tgataagaag	ogacttattt	ataatdatta	catatttttc	tattggaatg	attaagatto	4 20
16 1				gaaaggaggg			1,.,11
1 7				atttatgaaa			€, CPJ
171	aattatqtat	tatgataaga	aagggaggaa	gaaaaatgaa	toogaacaat	cgaagtgaac	$E_1 E_1 Q$
1 /	=			aggtgccaad			7. 0
1 7 .	tagoggaaac	tocaaatoca	ababtagaag	atttaaatta	taaagagtii	ttaagaatga	750
17.4	otycagataa	taatacggaa	goadtagata	gototadaad	aaaagatgto	attcaaaaag	544
1 70	gbatttdbgt	agtaggtgat	atostaggag	tagtaggttt	coogsttaggt	ggagogottg	ыÇÜ
1 76	tittogtittta	tacaaacttt	ttaaatasta	tttggddaag	tgaagacccg	tggaaggett	क्राहेस
1	ttatqqaaca	agtagaagca	ttgatggatc	agaaaatago	tgattatgda	aaaaataaag	1:0.55
1	otottgbaga	gttacagggc	ottoaaaata	atglogaaga	statgtgagt	gdattgagtt	1020
1,	patiggoaaaa	aaatootgtg	agttcacgaa	atocacatag	coaggggagg	ataagagago	1 1 4 . 1
1 : . :	tgittitatba	agcagaaagt	cattttogta	attomatgod	ttagtttgaa	atttotggat	1. (0)
1 1				aagotgoosa			1. 6.4
1:.	aagaogotoa	aatttatgga	gaagaatggg	gatacgaaaa	agaagatatt	gotgaatttt	1 5. 0
1	ataasagaba	actaaaactt	acgcaagaat	atactgacca	ttgtgtcaaa	tggtataatg	1 550
1 :	tit ggatit aga	taaattaaga	ggttcatctt	atgaatettg	ggtaaacttt	aaccgttatc	144:1
1	gcagagagat	gacattaada	gtattagatt	taattgoact	atttddattg	tatgatgttd	$1 \le (0.0)$
$1 - \epsilon$				taacaagaga			1560
1	toggagtoaa	caacettagg	ggetatggaa	daaddttata	taatatagaa	aattatatto	16.30
1 %				gaattdaatt			1600
18 4	gatattatgg	aaatgactot	ttdaattatt	ggtooggtaa	ttatgtttca	actagaccaa	1
100				cattotatgg			1800
191				totatagago			1-10
1 3.	aggiaiggee	gtoogotgta	tattcaggtg	ttacaaaagt	ggaatttagd	caatataatg	1
19.	atcaaacaga	tgaagbaagt	acacaaacgt	acgastsaaa	aagaaatgtt	ggogoggtoa	1 47 0
194	gotgggatto	tatugatdaa	ttgcctccag	aaadaadaga	tgaacctota	gaaaagggat	2 40
130	atagodatda	adtdaattat	gtaatgtgct	tittaatgda	gggtagtaga	ggaabaatbo	31.0
1 100	dagtigttaad	ttggadadat	aaaagtgtag	acttttttaa	catgattgat	togaaaaaaa	2160
1 -	ttadadaadt	toogttagta	aaggcatata	agttadaatd	tagtaettes	gttgtogcag	:
196	gtostaggtt	tabaqqaqqa	gatatoatto	aatgcacaga	aaatggaagt	goggosacta	.1.7 % 5.
19 •				asaaatstog			3.540
β¢π	otabatidtba	gataacattt	adactdagtt	tagacggggc	addatttaat	caatactatt	1.11
200				taacgtataa			14 × Q
.200				acttacaaat			2500
20.5				aatttattcc			$\mathcal{H}_{\mathcal{F}} \models C_{\mathcal{F}}$
.) Ü.,				taagcaaagg			jir 41.
205				dittitgaag			2700
200				tgtaatcatt			1111
.:0::	3 3			catttttata			.181.0
.:0:8				aaacgttcta			1188.0
.1011				tocatogagt			111440
.110				tgattggttg		gcgtaattta	,5(+()).
211	tgagggettt	tottttcato	aaaagccctc	gtgtatttct	ctgtaagctt		3050

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/834,271A TIME: 07:36:19

DATE: 08/05/2003

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt
Cutput Set: N:\CRF4\08052003\I834271A.raw

```
213 H2105 SEC ID No: 22
214 -0.11 - LENGTH: 17
215 3212 TYPE: EMA
216 C.13 · OF GANISM: Pacillus
31 F HAOF - SEQUENCE: 13
                                                                                        17
.:1:
          garettalig goetgea
225 (216 - SEQ TO NO: 23
ddi -011 - DENGTH: .2
dd. dd. L - TYFE: DHA
22 × 0.17 × OPGANISM: Pabillus
BB4 - 0400 - SEQUENCE: DB
                                                                                        22
Mill targing motitation to
327 K. 10 - SEO ID NO: 74
der - deli - Length: la
3339 KUTU TYFE: DNA
200 0.10 OBJANISM: Babillus
RNI 440: - SEQUENCE: 14
                                                                                        38
          gagetreatt trettataca aattatat
254 (21: SE, ID NO: .5
BEN SELLI - LENGTH: 185
down to 1. - Type: SMA
Had the ORGANISM: Madillus
day HIGH - SECUENCE: 13
                                                                                       (\tilde{\epsilon}, \tilde{\epsilon})
. . . . .
          gamestanga meetgeaate gattgtttga gaaaagaaga agaccataaa aataccttgt
          rtatostosa scagggtatt tittatgotg topagacigt cogpigigia aaaaatagga
                                                                                      120
. 1 -5 -
. . . 1
          aimandoqis istiqttatta tittactgat atqtaaaata taatttgtat aagaaaatgg
                                                                                      180
          a dintid
                                                                                       135
144 - 210 - SEQ 10 NO: 16
.4: -011 - LENGTH: 185
240 1.11 TYPE: DNA
347 -3313 - OFGANISM: Fabillus
NAS KAOO - SEQUENCE: 10
34 +
                                                                                       6.0
          pacettaung setqeaate qattqtttga qaaaaqaaqa agaccataaa aatacettgt
          ctatestess spaggitatt tittatgetg tecagactgt cegetgitgta aaaaaaagga
115.00
                                                                                      1. . . .
          ataaaggggg gttgacatta ttttactgat atgtataata taatttgtat aagaaaatgg
:5 j
                                                                                      180
          agoto
                                                                                      195
1354 -0.100 SEQ ID NO: 17
Mit -Will LENGTH: 185
23+ KIII - TYPE: DNA
. FT of 150 CFGANISM: Facillus
.15 K4000 SEQUENCE: 1/
                                                                                       ΕÜ
         - quinctiaaang dectgeaate gattgtttga gaaaagaaga agaccataaa aataccttgt
111
                                                                                      100
          cturearcag acagggraft tittatgetg tecagactgt cegetgigta aaaaatagga
atuaadedda ditqacatta ttttactgat atgtataata taatttgtat aagaaaatgg
                                                                                      180
                                                                                      165
          aquito
104 KI1CH SEQ II NO: 28
Def adding LENGTH: 53
200 KL12: TYPE: DNA
267 C 13: OKGANISM: Facillus
```

VERIFICATION SUMMARYDATE: 08/05/2003PATENT APPLICATION: US/09/834,271ATIME: 07:36:20

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt Output Set: N:\CRF4\08052003\1834271A.raw

L:12 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD

STATISTICS SUMMARY

PATENT APPLICATION: US/09/834,271A

DATE: 08/05/2003 TIME: 07:36:20

Input Set : N:\Crf3\RULE60\09834271A.RAW.txt
Output Set: N:\CRF4\08052003\I834271A.raw

Application Serial Number: US '09/834,271A

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 04-12-2001

Art Unit: OIPE

Software Application: FastSEQ3.0 Total Number of Sequences: 35 Total Nucleotides: 4542 Total Amino Acids: 0 Number of Errors: 0

Number of Warnings: 1 Number of Corrections: 0

MESSAGE SUMMARY

256 W: 1 (Invalid Numeric Header Field)